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L1 QUE EUKARYO? AND INITI? AND FACTO? AND KINAS?

FILE 'USPATFULL, ESBIOBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE, GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA, CABA, NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS' ENTERED AT 10:48:09 ON 14 NOV 2001

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- L4 9902 S L2 NOT L3
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- L7 1 S L5 AND 14790?
- L8 240 S L5 AND HUMAN? AND RECOMBIN?

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62 FILES IN THE FILE LIST IN STNINDEX

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=> s (eukaryo? and initi? and facto? and kinas?) or (eIF-2? and kinas?)

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39 FILE AGRICOLA

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          1 FILE NAPRALERT
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39 FILES HAVE ONE OR MORE ANSWERS, 62 FILES SEARCHED IN STNINDEX

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               ESBIOBASE
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7 FILES SEARCHED...

8 FILES SEARCHED... 13 FILES SEARCHED... 13541 EUKARYO? AND INITI? AND FACTO? AND KINAS? => s eIF-2? and kinas? TERM '2?' EXCEEDED TRUNCATION LIMITS - SEARCH ENDED 1 FILES SEARCHED... <-----User Break----> SEARCH ENDED BY USER You have entered a truncated stem which occurs in too many terms. Make the stem longer and try again. For example, if your original term was 'degr?' to search for variations and the abbreviation for 'degradation', you could replace it with the expression '(degrdn OR degrad?)'. If your search term was numeric, e.g., 'C>5', reduce the size of the range. => log hCOST IN U.S. DOLLARS SINCE FILE TOTAL SESSION ENTRY FULL ESTIMATED COST 35.01 76.56 SESSION WILL BE HELD FOR 60 MINUTES STN INTERNATIONAL SESSION SUSPENDED AT 10:54:03 ON 14 NOV 2001 Trying 3106016892...Open Welcome to STN International! Enter x: Welcome to STN International! Enter x:x LOGINID: LOGINID:ssspta1652dmr PASSWORD: * * * * * RECONNECTED TO STN INTERNATIONAL * * * * * SESSION RESUMED IN FILE 'USPATFULL, ESBIOBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE, GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA, CABA, NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS' AT 11:14:16 ON 14 NOV 2001 FILE 'USPATFULL' ENTERED AT 11:14:16 ON 14 NOV 2001 CA INDEXING COPYRIGHT (C) 2001 AMERICAN CHEMICAL SOCIETY (ACS) FILE 'ESBIOBASE' ENTERED AT 11:14:16 ON 14 NOV 2001 COPYRIGHT (C) 2001 Elsevier Science B.V., Amsterdam. All rights reserved. FILE 'BIOSIS' ENTERED AT 11:14:16 ON 14 NOV 2001 COPYRIGHT (C) 2001 BIOSIS(R) FILE 'SCISEARCH' ENTERED AT 11:14:16 ON 14 NOV 2001 COPYRIGHT (C) 2001 Institute for Scientific Information (ISI) (R) FILE 'CAPLUS' ENTERED AT 11:14:16 ON 14 NOV 2001 COPYRIGHT (C) 2001 AMERICAN CHEMICAL SOCIETY (ACS) FILE 'MEDLINE' ENTERED AT 11:14:16 ON 14 NOV 2001 FILE 'EMBASE' ENTERED AT 11:14:16 ON 14 NOV 2001 COPYRIGHT (C) 2001 Elsevier Science B.V. All rights reserved. FILE 'GENBANK' ENTERED AT 11:14:16 ON 14 NOV 2001 FILE 'BIOTECHNO' ENTERED AT 11:14:16 ON 14 NOV 2001 COPYRIGHT (C) 2001 Elsevier Science B.V., Amsterdam. All rights reserved. FILE 'LIFESCI' ENTERED AT 11:14:16 ON 14 NOV 2001 COPYRIGHT (C) 2001 Cambridge Scientific Abstracts (CSA) FILE 'CANCERLIT' ENTERED AT 11:14:16 ON 14 NOV 2001 FILE 'DGENE' ENTERED AT 11:14:16 ON 14 NOV 2001 COPYRIGHT (C) 2001 DERWENT INFORMATION LTD FILE 'PASCAL' ENTERED AT 11:14:16 ON 14 NOV 2001 COPYRIGHT (C) 2001 INIST-CNRS. All rights reserved. FILE 'TOXLIT' ENTERED AT 11:14:16 ON 14 NOV 2001 FILE 'AGRICOLA' ENTERED AT 11:14:16 ON 14 NOV 2001 FILE 'CABA' ENTERED AT 11:14:16 ON 14 NOV 2001

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18 FILES SEARCHED... 9902 L2 NOT L3 L4=> dup rem 12

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2528 DUP REM L3 (4761 DUPLICATES REMOVED) T.5

=> dup rem 12 DUPLICATE IS NOT AVAILABLE IN 'GENBANK, DGENE'. ANSWERS FROM THESE FILES WILL BE CONSIDERED UNIQUE PROCESSING IS APPROXIMATELY 5% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 9% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 14% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 24% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 30% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 36% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 42% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 42% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 48% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 54% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 76% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 77% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 77% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 85% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 77% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY 85% COMPLETE FOR L2 9% COMPLETE FOR L2 PROCESSING IS APPROXIMATELY

PROCESSING IS APPROXIMATELY 85% COMPLETE FOR L2

PROCESSING IS APPROXIMATELY 96% COMPLETE FOR L2

PROCESSING COMPLETED FOR L2 L6

10028 DUP REM L2 (3513 DUPLICATES REMOVED)

=> d hi

'HI' IS NOT A VALID FORMAT

In a multifile environment, a format can only be used if it is valid in at least one of the files. Refer to file specific help messages or the STNGUIDE file for information on formats available in individual files.

REENTER DISPLAY FORMAT FOR ALL FILES (FILEDEFAULT):end

(FILE 'HOME' ENTERED AT 09:52:28 ON 14 NOV 2001)

INDEX 'ADISALERTS, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, AQUASCI, BIOBUSINESS, BIOCOMMERCE, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CANCERLIT, CAPLUS, CEABA-VTB, CEN, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DRUGB, DRUGLAUNCH, DRUGMONOG2, ...' ENTERED AT 09:52:52 ON 14 NOV 2001

SEA (EUKARYO? AND INITI? AND FACTO? AND KINAS?) OR (EIF-2? AND

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1 FILE ADISALERTS
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- 39 FILE AGRICOLA
- 5 FILE AQUASCI
- 2 FILE BIOBUSINESS
- 1 FILE BIOCOMMERCE
- 912 FILE BIOSIS
 - 9 FILE BIOTECHABS
 - 9 FILE BIOTECHDS
- 501 FILE BIOTECHNO
- 25 FILE CABA
- 548 FILE CANCERLIT
 - 0* FILE CAPLUS
 - 1 FILE CEABA-VTB
 - 2 FILE CEN
- 23 FILE CONFSCI
- 14 FILE DDFU
- 160 FILE DGENE
 - 1 FILE DRUGNL
- 21 FILE DRUGU
- 14 FILE EMBAL
- 673 FILE EMBASE
- 795 FILE ESBIOBASE
 - 1 FILE FROSTI
 - 1 FILE FSTA
 - 0* FILE GENBANK
 - 0* FILE IFIPAT
- 11 FILE JICST-EPLUS
- 463 FILE LIFESCI
- 1256 FILE MEDLINE
 - 1 FILE NIOSHTIC
 - 0* FILE NTIS
 - 0* FILE PASCAL
 - 1 FILE PHIN
 - 0* FILE PROMT
 - 923 FILE SCISEARCH
 - 0* FILE TOXLIT
 - 0* FILE USPATFULL
 - 0* FILE WPIDS
 - 0* FILE WPINDEX
 - 1 FILE NAPRALERT
 - 0* FILE NLDB

SEA EUKARYO? AND INITI? AND FACTO? AND KINAS?

- 32 FILE AGRICOLA
- 4 FILE AQUASCI
- 2 FILE BIOBUSINESS
- 1 FILE BIOCOMMERCE
- 717 FILE BIOSIS
 - 9 FILE BIOTECHABS
 - 9 FILE BIOTECHDS
- 387 FILE BIOTECHNO
- 24 FILE CABA
- 210 FILE CANCERLIT

¹ FILE ADISALERTS

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2
                   FILE CEN
               6
                   FILE CONFSCI
                   FILE DDFU
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                   FILE DGENE
             129
                   FILE DRUGNL
              1
                   FILE DRUGU
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                   FILE EMBAL
              14
                   FILE EMBASE
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             746
                   FILE ESBIOBASE
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                   FILE FSTA
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                  FILE IFIPAT
              10
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             334
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                   FILE PROMT
               7
                   FILE SCISEARCH
              693
                   FILE TOXLIT
             116
            7669
                   FILE USPATFULL
              11
                   FILE WPIDS
              11
                   FILE WPINDEX
                  FILE NAPRALERT
               1
               19
                  FILE NLDB
                 QUE EUKARYO? AND INITI? AND FACTO? AND KINAS?
L1
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     CABA, NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS' ENTERED AT 10:48:09
     ON 14 NOV 2001
          13541 S EUKARYO? AND INITI? AND FACTO? AND KINAS?
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           7289 S EIF? AND KINAS?
L3
           9902 S L2 NOT L3
L4
           2528 DUP REM L3 (4761 DUPLICATES REMOVED)
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L6
          10028 DUP REM L2 (3513 DUPLICATES REMOVED)
=> s 15 and 14790?
   5 FILES SEARCHED...
  17 FILES SEARCHED...
             1 L5 AND 14790?
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    ANSWER 1 OF 1
                         Direct Submission
   TITLE (TI):
=> d 17
                           GENBANK.RTM. COPYRIGHT 2001
     ANSWER 1 OF 1
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                                      GenBank (R)
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GenBank ACC. NO. (GBN): Z71255
CAS REGISTRY NO. (RN):
                         175825-73-7
SEQUENCE LENGTH (SQL): 165536
                         DNA; linear
MOLECULE TYPE (CI):
DIVISION CODE (CI):
                         Plants, fungi, algae
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FILE CAPLUS

659

11 Oct 2000 DATE (DATE): S.cerevisiae chromosome XVI 165536 bp sequence, cen DEFINITION (DEF): rightwards. ARO7; ATPase; beta-transducin; CCL1; CDC21 homologue; KEYWORDS (ST): CDE1; CDE2; CDE3; centromere; CHL1; chorismate mutase; citrate synthase; coatomer zeta subunit; cyclin; delta element; DnaJ; DSS4; electro; transport flavoprotein; eukaryotic iniation factor 5; F1Fo-AT; synthase g subunit; gamma adaptin; GLN1; glutamine synthetase; Grave's disease; quanine-releasing protein; HAL1; halotolerance; histidine tRNA synthetase; HTS1; isocitrate lyase; LPT1; LTR; MAK3; MFS1; Mitochondrial carrier protein; N-acetyltransferase; NADPH-cytochrome P450 reductase; negative regulator; NHP6A; nonhistone chromosomal protein 6A; OSD1; phenylalanyl-tRN; synthetase; phosphotyrosine protein phosphatase; polyposis; protamine P1; protein kinase; protein transport; regulator; protein; ribosomal protein L37a; RNA polymerase I; ROX1; RPA2; SEC8; SMK1; spermidine synthase; TIF5; TKL1; transfer RNA-Gly; transfer RNA-Lys; transfer RNA-Phe; transfer RNA-Ser; transketolase; tropomyosin; ubiquitin-activating enzyme; vacuola; H(+)-ATPase 54 kda subunit; VMA13; YMC1; YME1; YTA11; zinc finge; protein baker's yeast. SOURCE: Saccharomyces cerevisiae ORGANISM (ORGN): Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes; Saccharomycetales; Saccharomycetaceae; Saccharomyces NUCLEIC ACID COUNT (NA): 52201 a 31306 c 31323 g 50706 t COMMENT: 1 to 25540 SC8132X Z48483 SC8132X Z48483 1 to 25540 SC9723 Z48951 100 to 17660 SC9531X Z49919 104 to 27166 SC9367 Z49274 10 to 42190 790 to 10339 (PCR product) Z68111 SC9499X Z49219 134 to 43776 The most significant matches using FASTA -o to a non-redundant database compiled from Swissprot 32 and PIR 46 and Wormpep9 are All CDS over 100 codons have been analysed. CDS that are largely or completely overlapped by a larger CDS have been omitted from this analysis. The more significant matches with motifs in the PROSITE database are also included but some of these may be fortuitous. The length in codons/amino-acids is given for each CDS as is the calculated codon adaptation index (CAI). tRNA genes and LTRs are also included. Notes: This sequence has been compiled from the following submissions of individual contigs. REFERENCE: (bases 1 to 165536) AUTHOR (AU): Badcock, K.; Bowman, S.; Churcher, C.M.; Pearson, D.; Rajandream, M.A.; Walsh, S.V.; Barrell, B.G. TITLE (TI): Direct Submission Submitted (19-APR-1996) Saccharomyces cerevisiae JOURNAL (SO): chromosome IV sequencing project, Sanger Centre, Hinxton Hall, Hinxton, Cambridge CB10 1RQ. E-mail: barrell@sanger.ac.uk FEATURES (FEAT): Feature Key Qualifier Location

SLQWARGTVYCTPAQLNELLGGVNRPASKYKESI IAEGSFSERSSDDNNANHPEHQYT KPTRKGRNSSYGVLRNVSRAVETRWDAVEDRFND YATAMGETMNEAVEQTGKDVMKGA LGF" /gene="YPR031W" 104601..104645 misc-feature /note="PS00678 Beta-transducin family Trp-Asp repeats signature" complement (107002..1086 /gene="HTS1" gene complement(107002..1086 /gene="HTS1" CDS 42) /note="YPR032C, HTS1 gene, len: 546, CAI: 0.30, SYH-YEAST P07263 histidyl-tRNA synthetase, contains PS00179 Aminoacyl-transfer RNA synthetases class-II signature 1 and PS00339 Aminoacyl-transfer RNA synthetases class-II signature 2" /codon-start=1 /product="Hts1p" /protein-id="CAA94983.1" /db-xref="GI:1314107" /db-xref="SWISS-PROT:P07263" /translation="MLSRSLNKVVTSIKSSSIIR

<---->

=> d his

(FILE 'HOME' ENTERED AT 09:52:28 ON 14 NOV 2001)

INDEX 'ADISALERTS, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, AQUASCI, BIOBUSINESS, BIOCOMMERCE, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CANCERLIT, CAPLUS, CEABA-VTB, CEN, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DRUGB, DRUGLAUNCH, DRUGMONOG2, ...' ENTERED AT 09:52:52 ON 14 NOV 2001

SEA (EUKARYO? AND INITI? AND FACTO? AND KINAS?) OR (EIF-2? AND

MSSATAAATSAPTANAANALKASK

LSGLFKKHGGVTIDTPVFELREIL

APKKGKLQVSLKTPKGTKDWADSDMVIREAIFST

AGKYGEDSKLIYNLEDQGGELCSLRYDLTVPFAR

1 FILE ADISALERTS 39 FILE AGRICOLA 5 FILE AQUASCI 2 FILE BIOBUSINESS FILE BIOCOMMERCE 1 912 FILE BIOSIS g FILE BIOTECHABS FILE BIOTECHDS G. 501 FILE BIOTECHNO FILE CABA 25 FILE CANCERLIT 548 0* FILE CAPLUS 1 FILE CEABA-VTB 2 FILE CEN 23 FILE CONFSCI 14 FILE DDFU 160 FILE DGENE 1 FILE DRUGNL 21 FILE DRUGU FILE EMBAL 14

FILE EMBASE

FILE ESBIOBASE

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795

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  0* FILE IFIPAT
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     FILE LIFESCI
1256
      FILE MEDLINE
      FILE NIOSHTIC
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  0* FILE NTIS
  0* FILE PASCAL
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  0* FILE USPATFULL
  0* FILE WPIDS
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      FILE NAPRALERT
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  0* FILE NLDB
   SEA EUKARYO? AND INITI? AND FACTO? AND KINAS?
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      FILE AQUASCI
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     FILE BIOTECHABS
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   9 FILE BIOTECHDS
     FILE BIOTECHNO
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     FILE EMBASE
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      FILE MEDLINE
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      FILE WPINDEX
  11
     FILE NAPRALERT
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  19 FILE NLDB
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L1

FILE 'USPATFULL, ESBIOBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE, GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA, CABA, NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS' ENTERED AT 10:48:09 ON 14 NOV 2001

13541 S EUKARYO? AND INITI? AND FACTO? AND KINAS? L2

7289 S EIF? AND KINAS? L3

9902 S L2 NOT L3 T.4

2528 DUP REM L3 (4761 DUPLICATES REMOVED) L5

10028 DUP REM L2 (3513 DUPLICATES REMOVED) T.6

1 S L5 AND 14790? L7

=> s 15 and human? and recombin?

3 FILES SEARCHED...

7 FILES SEARCHED...

11 FILES SEARCHED...

13 FILES SEARCHED...

240 L5 AND HUMAN? AND RECOMBIN?

=> s 16 and human? and recombin?

COMMAND INTERRUPTED

REENTER FILE 'USPATFULL, ESBIOBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE, GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA, CABA, NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS'

AND TRY AGAIN, OR ENTER '?' FOR MORE INFORMATION.

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=> s 16 and human? and recombin? THIS COMMAND NOT AVAILABLE IN THE CURRENT FILE Some commands only work in certain files. For example, the EXPAND command can only be used to look at the index in a file which has an index. Enter "HELP COMMANDS" at an arrow prompt (=>) for a list of commands which can be used in this file.

=> FIL F1-F24 COST IN U.S. DOLLARS

TOTAL SINCE FILE ENTRY SESSION 125.75 167.30

FULL ESTIMATED COST

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=> d his

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FILE EMBASE 520 746 FILE ESBIOBASE 1 FILE FROSTI 1 FILE FSTA 491 FILE GENBANK FILE IFIPAT FILE JICST-EPLUS 10 334 FILE LIFESCI FILE MEDLINE 620 1 FILE NIOSHTIC 116 FILE PASCAL FILE PHIN 1 FILE PROMT 7 693 FILE SCISEARCH 116 FILE TOXLIT FILE USPATFULL 7669 FILE WPIDS 11 FILE WPINDEX 11 FILE NAPRALERT FILE NLDB QUE EUKARYO? AND INITI? AND FACTO? AND KINAS?

FILE 'USPATFULL, ESBIOBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE, GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA, CABA, NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS' ENTERED AT 10:48:09 ON 14 NOV 2001

L2 13541 S EUKARYO? AND INITI? AND FACTO? AND KINAS?

L3 7289 S EIF? AND KINAS?

L4 9902 S L2 NOT L3

L1

L7

L5 2528 DUP REM L3 (4761 DUPLICATES REMOVED)

L6 10028 DUP REM L2 (3513 DUPLICATES REMOVED)

1 S L5 AND 14790?

L8 240 S L5 AND HUMAN? AND RECOMBIN?

FILE 'USPATFULL, ESBIOBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE, GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA, CABA, NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS' ENTERED AT 12:28:03 ON 14 NOV 2001

=> s 16 and human? and recombin?

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REENTER FILE 'USPATFULL, ESBIOBASE, BIOSIS, SCISEARCH, CAPLUS, MEDLINE, EMBASE, GENBANK, BIOTECHNO, LIFESCI, CANCERLIT, DGENE, PASCAL, TOXLIT, AGRICOLA, CABA, NLDB, DRUGU, EMBAL, WPIDS, IFIPAT, JICST-EPLUS'

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=> s 18 and alph?
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=> FIL F1-F24 COST IN U.S. DOLLARS

FULL ESTIMATED COST

SINCE FILE TOTAL
ENTRY SESSION
27.48 194.78

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hyper-thermophilic crenarchaeon, Aeropyrum pernix K1

Direct Submission TITLE (TI):

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BIOBUSINESS, BIOCOMMERCE, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CANCERLIT, CAPLUS, CEABA-VTB, CEN, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DRUGB, DRUGLAUNCH, DRUGMONOG2, ...' ENTERED AT 09:52:52 ON 14 NOV 2001

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177 S L8 AND ALPH? T.9

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